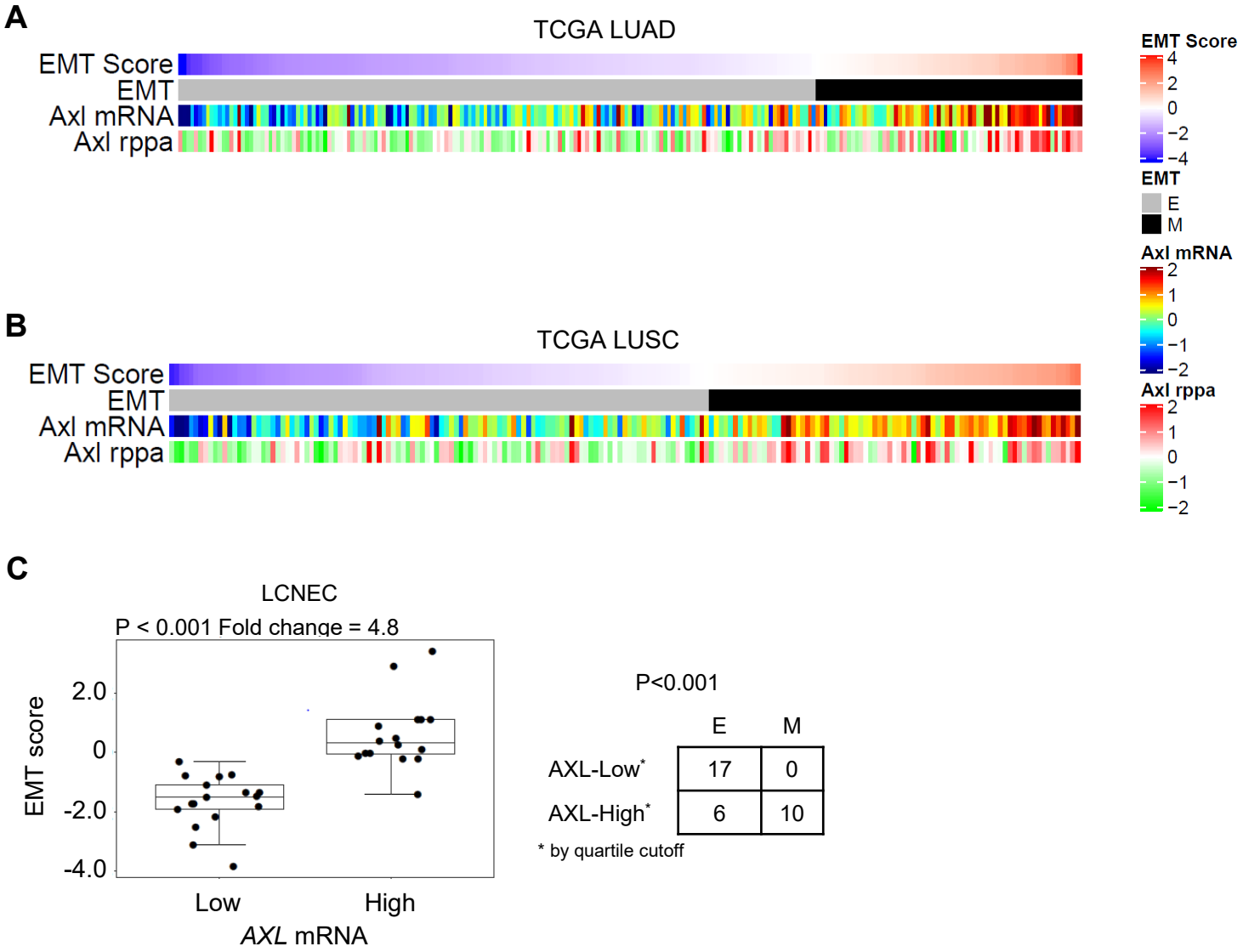

Supplementary Fig. S1. A subset of treatment-naïve NSCLC and LCNEC tumors expresses high AXL levels and EMT scores. **A** and **B.** Heatmap depicts the range of EMT scores, *AXL* mRNA and AXL protein expression of treatment-naïve lung adenocarcinomas (TCGA LUAD) (**A**) and lung squamous cell carcinomas (TCGA LUSC) (**B**). (**C**) Comparison of EMT scores and frequency of E and M tumors between LCNEC tumors with high and low *AXL* mRNA levels, determined by quartile cutoff.